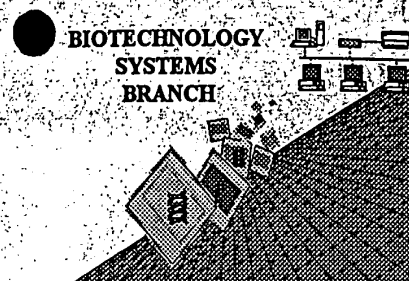


Wilson

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number: 09/205,096
Art Unit / Team No: 1633
Date Processed by STIC: 2/16/2000

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,

2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/205,096

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 ☐ Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 ☐ Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 ☐ Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 ☐ Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 ☐ Variable Length Sequence(s) ☐ contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 ☐ PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) ☐. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence.
- 8 ☐ Skipped Sequences (OLD RULES) Sequence(s) ☐ missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 ☐ Skipped Sequences (NEW RULES) Sequence(s) ☐ missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 ☐ Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 ☐ Use of <213>Organism (NEW RULES) Sequence(s) ☐ are missing this mandatory field or its response.
- 12 ☒ Use of <220>Feature (NEW RULES) Sequence(s) ☐ are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 ☐ PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory number identifiers and responses (as indicated on raw sequence listing).
Instead, please use "File Manager" or any other means to copy file to floppy disk.
AKS-Biotechnology Systems Branch- 5/15/99

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/205,096DATE: 02/16/2000
TIME: 11:05:46

Input Set: I205096.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

Does Not Comply
Corrected Diskette Needed

1 <110> APPLICANT: Drachman, Daniel B.
2 <120> TITLE OF INVENTION: TARGETING ANTIGEN-SPECIFIC T CELLS FOR
3 SPECIFIC IMMUNOTHERAPY OF AUTOIMMUNE DISEASE
4 <130> FILE REFERENCE: 01107.77737
5 <140> CURRENT APPLICATION NUMBER: US/09/205,096
6 <141> CURRENT FILING DATE: 1998-12-03
7 <150> EARLIER APPLICATION NUMBER: 60/067547
8 <151> EARLIER FILING DATE: 1998-12-03
9 <160> NUMBER OF SEQ ID NOS: 6
10 <170> SOFTWARE: FastSEQ for Windows Version 3.0
11 <210> SEQ ID NO 1
12 <211> LENGTH: 39
13 <212> TYPE: DNA
14 <213> ORGANISM: Artificial Sequence
15 <220> FEATURE:
16 <223> OTHER INFORMATION: designed primer
17 <400> SEQUENCE: 1
18 catatggatc caagcttatg attctgtgca gttattggc 39
19 <210> SEQ ID NO 2
20 <211> LENGTH: 37
21 <212> TYPE: DNA
22 <213> ORGANISM: Artificial Sequence
23 <400> SEQUENCE: 2
24 aggcctcgag aattcaatac gctgcatgat aaaatgg 37
25 <210> SEQ ID NO 3
26 <211> LENGTH: 26
27 <212> TYPE: DNA
28 <213> ORGANISM: Artificial Sequence
29 <400> SEQUENCE: 3
30 ggggaagctt atggcggccc ccggcg 26
31 <210> SEQ ID NO 4
32 <211> LENGTH: 28
33 <212> TYPE: DNA
34 <213> ORGANISM: Artificial Sequence
35 <400> SEQUENCE: 4
36 atgcaagctt tagatcctca aagagtgc 28
37 <210> SEQ ID NO 5
38 <211> LENGTH: 31
39 <212> TYPE: DNA
40 <213> ORGANISM: Artificial Sequence
41 <400> SEQUENCE: 5
42 gggggaattc ttgatcccca ttgctgtggg c 31
43 <210> SEQ ID NO 6
44 <211> LENGTH: 31

see item 12 on Error Summary sheet

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/205,096DATE: 02/16/2000
TIME: 11:05:46

Input Set: I205096.RAW

45 <212> TYPE: DNA
46 <213> ORGANISM: Artificial Sequence
47 <400> SEQUENCE: 6
48 aaaactcgag ctagatgggc tgatagccgg c

31

PAGE: 3

VERIFICATION SUMMARY
PATENT APPLICATION US/09/205,096

DATE: 02/16/2000
TIME: 11:05:46

Input Set: I205096.RAW

Line ? Error/Warning

Original Text
